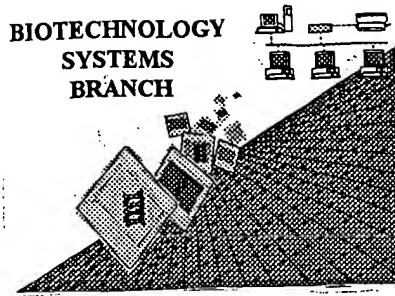


Tay

BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/926,163
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FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

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<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom, including:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission

User Manual - ePAVE)

2. U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202

3. Hand Carry directly to:

U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202

Or

U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,
2011 South Clark Place, Arlington, VA 22202

4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office,
Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

09/926/63

SPECIMEN SEQUENCE LISTING

Does Not Comply
Corrected Diskette Needed

<110> Fujisawa Pharmaceutical Co., Ltd.
<120> Sorbitol Dehydrogenase, Gene Coding Thereof and Use Thereof
<130> 09348
<150> JP 11-72810
<151> 1999-03-17
<150> JP 11-224679
<151> 1999-08-06
<160> 20
<210> 1
<211> 485
<212> PRT
<213> Gluconobacter oxydans

<400> 1
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@1 @ @ @ @ @ @ @ 5 @ @ @ @ @ @ @ @ 10 @ @ @ @ @ @ @ @ 15
Pro Pro Tyr Asp Ile Asp Gly Ile Lys Pro Gly Ile Val His Phe Gly
@ @ @ @ @ @ @ 20 @ @ @ @ @ @ @ @ 25 @ @ @ @ @ @ @ @ 30

(sample 1) submitted file)

delete
these
non-ASCII
characters
throughout
file, and
re-align amino
acid numbers. DO
NOT use TAB
codes between
amino acid numbers.

FYI →

Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.